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Sequence Listing was accepted.
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217-9197 (toll free).
Reviewer: Anne Corrigan
Timestamp: [year=2009; month=4; day=18; hr=17; min=48; sec=44; ms=450; ]
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Application No: 10562486   Version No: 3.0

Input Set:

Output Set:

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Finished: 2009-04-03 09:51:19.129
Elapsed: 0 hr(s) 0 min(s) 1 sec(s) 648 ms
Total Warnings: 21
Total Errors: 0
No. of SeqIDs Defined: 22
Actual SeqID Count: 22

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W 213	Artificial or Unknown found in <213> in SEQ ID (21)

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        Input Set:


       Output Set:



          Started:   2009-04-03 09:51:17.481
         Finished:   2009-04-03 09:51:19.129
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   Total Warnings:   21
     Total Errors:   0
No. of SeqIDs Defined:   22
 Actual SeqID Count:   22
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Error code   Error Description

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<110> Sugiyama, Haruo

<120> Method of selecting patients suitable for WT1 vaccine

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<141> 2005-12-27

<150> PCT/JP2004/009378
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<150> JP 2003-184436
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Gln Trp Ala Pro Val Leu Asp Phe Ala Pro Pro Gly Ala Ser Ala Tyr
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Gly Ser Leu Gly Gly Pro Ala Pro Pro Pro Ala Pro Pro Pro Pro Pro
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Pro Pro Pro Pro His Ser Phe Ile Lys Gln Glu Pro Ser Trp Gly Gly
 65                  70                  75                  80

Ala Glu Pro His Glu Glu Gln Cys Leu Ser Ala Phe Thr Val His Phe
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Ser Gly Gln Phe Thr Gly Thr Ala Gly Ala Cys Arg Tyr Gly Pro Phe
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Gly Pro Pro Pro Pro Ser Gln Ala Ser Ser Gly Gln Ala Arg Met Phe
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Pro Asn Ala Pro Tyr Leu Pro Ser Cys Leu Glu Ser Gln Pro Ala Ile
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Gly His Thr Pro Ser His His Ala Ala Gln Phe Pro Asn His Ser Phe
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Lys His Glu Asp Pro Met Gly Gln Gln Gly Ser Leu Gly Glu Gln Gln
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Tyr Ser Val Pro Pro Pro Val Tyr Gly Cys His Thr Pro Thr Asp Ser
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Cys Thr Gly Ser Gln Ala Leu Leu Leu Arg Thr Pro Tyr Ser Ser Asp
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Asn Leu Tyr Gln Met Thr Ser Gln Leu Glu Cys Met Thr Trp Asn Gln
225   230   235   240

Met Asn Leu Gly Ala Thr Leu Lys Gly Val Ala Ala Gly Ser Ser Ser
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Ser Val Lys Trp Thr Glu Gly Gln Ser Asn His Ser Thr Gly Tyr Glu
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His Thr His Gly Val Phe Arg Gly Ile Gln Asp Val Arg Arg Val Pro
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Gly Val Ala Pro Thr Leu Val Arg Ser Ala Ser Glu Thr Ser Glu Lys
305   310   315   320

Arg Pro Phe Met Cys Ala Tyr Pro Gly Cys Asn Lys Arg Tyr Phe Lys
                  325   330   335

Leu Ser His Leu Gln Met His Ser Arg Lys His Thr Gly Glu Lys Pro
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Tyr Gln Cys Asp Phe Lys Asp Cys Glu Arg Arg Phe Ser Arg Ser Asp
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Gln Leu Lys Arg His Gln Arg Arg His Thr Gly Val Lys Pro Phe Gln
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Cys Lys Thr Cys Gln Arg Lys Phe Ser Arg Ser Asp His Leu Lys Thr
385   390   395   400

His Thr Arg Thr His Thr Gly Lys Thr Ser Glu Lys Pro Phe Ser Cys
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